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This listing of the claims will replace all prior versions and listings of claims in the application:

Listing of the claims:

Claim 1 (canceled)

Claim 2 (previously amended): A method of diagnosing observations. Claim 2 (previously amended): A method of diagnosing comprising:

- obtaining a DNA or RNA sample from a patient; and
- screening the DNA or RNA sample with an oligonucleotide probe to detect a hMLH1 mutant 1, a hMSH2 mutant 1, a hMSH2 mutant 2, or a hMSH2 mutant 3, wherein binding of the oligonucleotide probe to the DNA or RNA sample is indicative of the presence of the hMLH1 mutant 1, the hMSH2 mutant 1, the hMSH2 mutant 2, or the hMSH2 mutant 3 and hereditary non-polyposis colorectal cancer.

Claim 3 (previously amended): A method for predicting susceptibility of a patient to developing hereditary nonpolyposis colorectal cancer comprising:

- obtaining a DNA or RNA sample from a patient; and
- screening the DNA or RNA sample with an (b) oligonucleotide probe to detect a hMLH1 mutant 1, a hMSH2 mutant 1, a hMSH2 mutant 2, or a hMSH2 mutant 3, wherein binding of the oligonucleotide probe to the DNA or RNA